RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number:	10/539,217
Source:	PCT
Date Processed by STIC:	06/09/2006
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ENTERED



PCT

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PATENT APPLICATION: US/10/539,217
                                                             TIME: 15:08:01
                     Input Set : A:\PTO.RJ.txt
                     Output Set: N:\CRF4\06092006\J539217.raw
      3 <110> APPLICANT: The Regents of the University of California
             Shokat, Kevan M.
             Knight, Zachary
      7 <120> TITLE OF INVENTION: Chemo-Enzymatic Process For Proteome-Wide Mapping of Post-
Translational
             Modification
     10 <130> FILE REFERENCE: 18062G-005410US
                                                                  CP96-7)
     12 <140> CURRENT APPLICATION NUMBER: US 10/539,217
     13 <141> CURRENT FILING DATE: 2005-06-17
     15 <150> PRIOR APPLICATION NUMBER: PCT/ US2003/041118
     16 <151> PRIOR FILING DATE: 2003-12-18
     18 <150> PRIOR APPLICATION NUMBER: US 60/434,696
     19 <151> PRIOR FILING DATE: 2002-12-18
     21 <160> NUMBER OF SEQ ID NOS: 57
     23 <170> SOFTWARE: PatentIn version 3.1
     25 <210> SEQ ID NO: 1
    26 <211> LENGTH: 7
     27 <212> TYPE: PRT
     28 <213> ORGANISM: Artificial Sequence
     30 <220> FEATURE:
     31 <223> OTHER INFORMATION: Peptide containing a cysteic acid residue
     33 <220> FEATURE:
     34 <221> NAME/KEY: MISC FEATURE
     35 <222> LOCATION: (5)..(5)
     36 <223> OTHER INFORMATION: Xaa = cysteic acid
    39 <400> SEQUENCE: 1
W--> 41 Leu Arg Arg Ala Xaa Leu Gly
     42 1
     45 <210> SEQ ID NO: 2
     46 <211> LENGTH: 4
     47 <212> TYPE: PRT
     48 <213> ORGANISM: Artificial Sequence
     50 <220> FEATURE:
    51 <223> OTHER INFORMATION: Digested peptide fragment after cleavage with peptidyl-
aspartate
    52
             metalloendopeptidase (Asp-N)
    54 <400> SEQUENCE: 2
    56 Leu Arg Arg Ala
    57 1
    60 <210> SEQ ID NO: 3
    61 <211> LENGTH: 9
    62 <212> TYPE: PRT
    63 <213> ORGANISM: Artificial Sequence
    65 <220> FEATURE:
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66 <223> OTHER INFORMATION: Undigested betamethyl aminoethylcysteine modified peptide

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     74 <220> FEATURE:
     75 <221> NAME/KEY: MISC_FEATURE
     76 <222> LOCATION: (8)..(8)
     77 <223> OTHER INFORMATION: Xaa = nitrotyrosine
     80 <400> SEQUENCE: 3
W--> 82 Glx Phe Arg Pro Xaa Gly Phe Xaa Glu
     83 1
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     87 <211> LENGTH: 9
     88 <212> TYPE: PRT
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     91 <220> FEATURE:
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     96 <222> LOCATION: (5)..(5)
     97 <223> OTHER INFORMATION: Xaa = phosphotyrosine
     100 <220> FEATURE:
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     103 <223> OTHER INFORMATION: Xaa = nitrotyrosine
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W--> 108 Glx Phe Arg Pro Xaa Gly Phe Xaa Glu
     109 1
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     121 <221> NAME/KEY: MISC FEATURE
     122 <222> LOCATION: (6)..(6)
     123 <223> OTHER INFORMATION: Xaa = aminoethylcysteine (K*)
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W--> 128 Asn Lys Pro Pro Arg Xaa Pro Val Val Glu Leu Ser Lys
     129 1
     132 <210> SEQ ID NO: 6
     133 <211> LENGTH: 15
     134 <212> TYPE: PRT
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     137 <220> FEATURE:
     138 <223> OTHER INFORMATION: Peptide containing a phosphoserine
     140 <220> FEATURE:
     141 <221> NAME/KEY: MISC FEATURE
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PATENT APPLICATION: US/10/539,217 TIME: 15:08:01 Input Set : A:\PTO.RJ.txt Output Set: N:\CRF4\06092006\J539217.raw 142 <222> LOCATION: (7)..(7) 143 <223> OTHER INFORMATION: Xaa = phosphoserine 146 <400> SEQUENCE: 6 W--> 148 Asn Lys Lys Pro Pro Arg Xaa Ser Pro Val Val Glu Leu Ser Lys 149 1 10 152 <210> SEO ID NO: 7 153 <211> LENGTH: 13 154 <212> TYPE: PRT 155 <213> ORGANISM: Artificial Sequence 157 <220> FEATURE: 158 <223> OTHER INFORMATION: unmodified peptide 160 <400> SEQUENCE: 7 162 Asn Lys Pro Pro Arg Ser Pro Val Val Glu Leu Ser Lys 163 1 166 <210> SEQ ID NO: 8 167 <211> LENGTH: 23 168 <212> TYPE: PRT 169 <213> ORGANISM: Artificial Sequence 171 <220> FEATURE: 172 <223> OTHER INFORMATION: Guanidinated MARCKS peptide 174 <220> FEATURE: 175 <221> NAME/KEY: MISC FEATURE 176 <222> LOCATION: (1)..(5) 177 <223> OTHER INFORMATION: Xaa = guanidinylysine 180 <220> FEATURE: 181 <221> NAME/KEY: MISC_FEATURE 182 <222> LOCATION: (8)..(8) 183 <223> OTHER INFORMATION: Xaa = aminoethylcysteine (K*) 186 <220> FEATURE: 187 <221> NAME/KEY: MISC FEATURE 188 <222> LOCATION: (10)..(12) 189 <223> OTHER INFORMATION: Xaa = guanidinylysine 192 <220> FEATURE: 193 <221> NAME/KEY: MISC FEATURE 194 <222> LOCATION: (14)..(14) 195 <223> OTHER INFORMATION: Xaa = aminoethylcysteine (K*) 198 <220> FEATURE: 199 <221> NAME/KEY: MISC FEATURE 200 <222> LOCATION: (19)..(20) 201 <223> OTHER INFORMATION: Xaa = guanidinylysine 204 <220> FEATURE: 205 <221> NAME/KEY: MISC FEATURE 206 <222> LOCATION: (22)..(23) 207 <223> OTHER INFORMATION: Xaa = guanidinylysine 210 <400> SEQUENCE: 8 W--> 212 Xaa Xaa Xaa Xaa Xaa Arg Phe Xaa Phe Xaa Xaa Aaa Phe Xaa Leu Ser 213 1 10 216 Gly Phe Xaa Xaa Asn Xaa Xaa 217 20

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PATENT APPLICATION: US/10/539,217
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     220 <210> SEQ ID NO: 9
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Lys-
               C (Lys-C)
     229 <220> FEATURE:
     230 <221> NAME/KEY: MISC FEATURE
     231 <222> LOCATION: (8)..(8)
     232 <223> OTHER INFORMATION: Xaa = aminoethylcysteine (K*)
     235 <400> SEQUENCE: 9
W--> 237 Lys Lys Lys Lys Arg Phe Xaa
     238 1
     241 <210> SEO ID NO: 10
     242 <211> LENGTH: 12
     243 <212> TYPE: PRT
     244 <213> ORGANISM: Artificial Sequence
     246 <220> FEATURE:
     247 <223> OTHER INFORMATION: Acetylated MARCKS peptide after cleavage with endoproteinase
Lys-
     248
               C (Lys-C)
     250 <220> FEATURE:
     251 <221> NAME/KEY: MISC FEATURE
     252 <222> LOCATION: (8)..(8)
     253 <223> OTHER INFORMATION: Xaa - aminoethylcysteine (K*)
     256 <220> FEATURE:
     257 <221> NAME/KEY: MISC FEATURE
     258 <222> LOCATION: (12)..(12)
     259 <223> OTHER INFORMATION: Xaa - aminoethylcysteine (K*)
     262 <400> SEQUENCE: 10
W--> 264 Lys Lys Lys Lys Arg Phe Xaa Phe Lys Lys Xaa
     268 <210> SEQ ID NO: 11
     269 <211> LENGTH: 19
     270 <212> TYPE: PRT
     271 <213> ORGANISM: Artificial Sequence
     273 <220> FEATURE:
     274 <223> OTHER INFORMATION: Acetylated MARCKS peptide after cleavage with endoproteinase
Lys-
     275
              C (Lys-C)
     277 <220> FEATURE:
     278 <221> NAME/KEY: MISC_FEATURE
     279 <222> LOCATION: (8)..(8)
     280 <223> OTHER INFORMATION: Xaa = aminoethylcysteine
     283 <220> FEATURE:
     284 <221> NAME/KEY: MISC FEATURE
     285 <222> LOCATION: (12)..(12)
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     289 <220> FEATURE:
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RAW SEQUENCE LISTING
                                                              DATE: 06/09/2006
                     PATENT APPLICATION: US/10/539,217
                                                             TIME: 15:08:01
                     Input Set : A:\PTO.RJ.txt
                     Output Set: N:\CRF4\06092006\J539217.raw
     291 <222> LOCATION: (19)..(19)
     292 <223> OTHER INFORMATION: Xaa = aminoethylcysteine
     295 <400> SEQUENCE: 11
W--> 297 Lys Lys Lys Lys Arg Phe Xaa Phe Lys Xaa Phe Lys Leu Ser
     298 1
                                              10
     301 Gly Phe Xaa
     305 <210> SEQ ID NO: 12
     306 <211> LENGTH: 25
     307 <212> TYPE: PRT
     308 <213> ORGANISM: Artificial Sequence
     310 <220> FEATURE:
     311 <223> OTHER INFORMATION: Acetylated MARCKS peptide after cleavage with endoproteinase
Lys-
     312
              C (Lys-C)
     314 <220> FEATURE:
     315 <221> NAME/KEY: MISC FEATURE
     316 <222> LOCATION: (8)..(8)
     317 <223> OTHER INFORMATION: Xaa = aminoethylcysteine
     320 <220> FEATURE:
     321 <221> NAME/KEY: MISC FEATURE
     322 <222> LOCATION: (12)..(12)
     323 <223> OTHER INFORMATION: Xaa = aminoethylcysteine
     326 <220> FEATURE:
     327 <221> NAME/KEY: MISC_FEATURE
     328 <222> LOCATION: (19)..(19)
     329 <223> OTHER INFORMATION: Xaa = aminoethylcysteine
     332 <400> SEQUENCE: 12
W--> 334 Lys Lys Lys Lys Arg Phe Xaa Phe Lys Lys Xaa Phe Lys Leu Ser
                         5
     338 Gly Phe Xaa Phe Lys Lys Asn Lys Lys
     339
                     20
     342 <210> SEQ ID NO: 13
     343 <211> LENGTH: 8
     344 <212> TYPE: PRT
     345 <213> ORGANISM: Artificial Sequence
     347 <220> FEATURE:
     348 <223> OTHER INFORMATION: Acetylated MARCKS peptide after cleavage with endoproteinase
Lys-
     349
              C (Lys-C)
     351 <220> FEATURE:
     352 <221> NAME/KEY: MISC FEATURE
     353 <222> LOCATION: (1)..(1)
    354 <223> OTHER INFORMATION: Xaa = aminoethylcysteine
     357 <220> FEATURE:
    358 <221> NAME/KEY: MISC FEATURE
    359 <222> LOCATION: (8)..(8)
    360 <223> OTHER INFORMATION: Xaa = aminoethylcysteine
    363 <400> SEQUENCE: 13
W--> 365 Xaa Phe Lys Leu Ser Gly Phe Xaa
    366 1
    369 <210> SEQ ID NO: 14
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Input Set : A:\PTO.RJ.txt

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Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

```
Seq#:1; Xaa Pos. 5
Seq#:3; Xaa Pos. 5,8
Seq#:4; Xaa Pos. 5,8
Seg#:5; Xaa Pos. 6
Seq#:6; Xaa Pos. 7
Seq#:8; Xaa Pos. 1,2,3,4,5,8,10,11,12,14,19,20,22,23
Seq#:9; Xaa Pos. 8
Seq#:10; Xaa Pos. 8,12
Seq#:11; Xaa Pos. 8,12,19
Seg#:12; Xaa Pos. 8,12,19
Seq#:13; Xaa Pos. 1,8
Seq#:14; Xaa Pos. 1
Seq#:15; Xaa Pos. 1,5,12
Seq#:16; Xaa Pos. 7
Seq#:17; Xaa Pos. 1,5,12
Seq#:18; Xaa Pos. 8
Seq#:19; Xaa Pos. 15
Seq#:20; Xaa Pos. 6
Seq#:21; Xaa Pos. 4
Seq#:22; Xaa Pos. 5
Seq#:23; Xaa Pos. 5
Seq#:24; Xaa Pos. 5
Seq#:25; Xaa Pos. 3
Seq#:26; Xaa Pos. 4,6,12
Seq#:27; Xaa Pos. 1,3,9
Seq#:28; Xaa Pos. 1,7
Seq#:29; Xaa Pos. 10
Seq#:30; Xaa Pos. 10
Seq#:31; Xaa Pos. 6
Seq#:32; Xaa Pos. 1
Seg#:33; Xaa Pos. 15,17,18,19
Seg#:34; Xaa Pos. 15,17,18,19
Seq#:35; Xaa Pos. 15,17,18
Seq#:36; Xaa Pos. 15,17
Seq#:37; Xaa Pos. 15
Seq#:38; Xaa Pos. 3
Seq#:39; Xaa Pos. 1
Seq#:40; Xaa Pos. 1,8,12,19
Seq#:41; Xaa Pos. 1,8,12,19
Seq#:42; Xaa Pos. 1,8,12
Seg#:43; Xaa Pos. 1,8
Seq#:44; Xaa Pos. 4,11
Seq#:45; Xaa Pos. 4,11
Seq#:46; Xaa Pos. 7
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Seq#:47; Xaa Pos. 1
Seq#:49; Xaa Pos. 5
Seq#:50; Xaa Pos. 3
Seq#:51; Xaa Pos. 1
Seq#:52; Xaa Pos. 6
Seq#:53; Xaa Pos. 3,6
Seq#:54; Xaa Pos. 6

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L:41 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:0
L:82 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:0
L:108 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4 after pos.:0
L:128 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5 after pos.:0
L:148 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6 after pos.:0
L:212 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8 after pos.:0
M:341 Repeated in SeqNo=8
L:237 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9 after pos.:0
L:264 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10 after pos.:0
L:297 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11 after pos.:0
M:341 Repeated in SeqNo=11
L:334 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:12 after pos.:0
M:341 Repeated in SeqNo=12
L:365 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13 after pos.:0
L:386 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:14 after pos.:0
L:419 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15 after pos.:0
L:440 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:16 after pos.:0
L:473 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:17 after pos.:0
L:497 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:18 after pos.:0
L:517 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:19 after pos.:0
L:541 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:20 after pos.:0
L:565 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:21 after pos.:0
L:585 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:22 after pos.:0
L:605 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:23 after pos.:0
L:625 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:24 after pos.:0
L:645 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:25 after pos.:0
L:678 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:26 after pos.:0
L:711 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:27 after pos.:0
L:738 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:28 after pos.:0
L:759 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:29 after pos.:0
L:780 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:30 after pos.:0
L:801 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:31 after pos.:0
L:822 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:32 after pos.:0
L:849 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:33 after pos.:0
M:341 Repeated in SeqNo=33
L:880 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:34 after pos.:0
M:341 Repeated in SeqNo=34
L:911 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:35 after pos.:0
M:341 Repeated in SeqNo=35
L:942 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:36 after pos.:0
M:341 Repeated in SeqNo=36
L:967 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:37 after pos.:0
L:988 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:38 after pos.:0
L:1009 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:39 after pos.:0
L:1048 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:40 after pos.:0
M:341 Repeated in SeqNo=40
L:1091 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:41 after pos.:0
M:341 Repeated in SeqNo=41
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L:1128 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:42 after pos.:0 L:1155 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:43 after pos.:0